AMENDMENTS TO THE CLAIMS

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- (WITHDRAWN-CURRENTLY AMENDED): A method <u>for detection of Yersinia</u> <u>pestis in a sample</u> comprising:
 - (i) providing a sample;
- (ii) forming a mixture by adding the sample to a solution containing at least one series of nucleotide sequences having a forward primer, a reverse primer and a hybridization probe selected from the group consisting of SEQ ID NOs:1, 2, 3; 5, 6, 7; 9, 10, 11; 13, 14, 15;17, 18, 19; 21, 22, 23; under conditions suitable for isolating genomic DNA for amplification using PCR and under conditions suitable for hybridization with said at least one series of nucleotide sequences; and
- (iii) subjecting the mixture to PCR using a PCR assay to detect the composition of claim 8 in the sample, wherein the detection of SEQ ID NO:4 and SEQ ID NO:8 in the sample indicates the presence of Yersinia pestis in the sample.
- (WITHDRAWN): The method of Claim 4 wherein said PCR comprises standard PCR.
- (WITHDRAWN): The method of Claim 5, wherein said PCR comprises fluorogenic
 nuclease PCR assay.
- (WITHDRAWN-CURRENTLY AMENDED): A method comprising The method of claim 4, wherein
 - (i) providing a sample;
- (ii) forming a mixture by adding the sample to a solution containing at least one series of nucleotide sequences having a forward primer, a reverse primer and a hybridization probe selected from the group consisting of SEQ ID NOS 1, 2, 3, 5, 6, 7; 9, 10, 11; 13, 14, 15; 17, 18, 19; 21, 22, 23;
- said assay is performed using a first forward primer consisting of SEQ ID NO:1, a first reverse primer consisting of SEQ ID NO:2, and a first hybridization probe consisting of SEQ ID

NO:3 for detection of SEQ ID NO:4 and using a second forward primer consisting of SEQ ID NO:5, a second reverse primer consisting of SEQ ID NO:6, and a second hybridization probe consisting of SEQ ID NO:7 for detection of SEQ ID NO:8.;

under conditions suitable for isolating genomic DNA for amplification using PCR and under conditions suitable for hybridization with said at least one series of nucleotide sequences; and

- 8. (CURRENTLY AMENDED) A composition comprising a first isolated polynucleotide and a second isolated polynucleotide, wherein the first isolated polynucleotide emprises-consists of SEQ ID NO:4 or a full-length complement thereof and the second polynucleotide emprises-consists of SEQ ID NO:8 or a full-length complement thereof.
- 9. (CURRENTLY AMENDED) The composition of claim 8, comprising at least one further isolated polynucleotide emprising consisting of a nucleic acid sequence selected from the group consisting of SEQ ID NOS: 12, 16, 20, and 24 or a full-length complement thereof.
- 10. (CURRENTLY AMENDED) The composition of claim 9, comprising six isolated polynucleotides each eomprising consisting of one of SEQ ID NOS: 4, 8, 12, 16, 20, and 24 or full-length complements thereof.
- 11. (CURRENTLY AMENDED) A set of oligonucleotides comprising (a) a polynucleotide fragment of each of the isolated polynucleotides of the composition of claim 8, wherein said fragments are 42 to 50 19 to 33 nucleotides in length, or (b) full-length complements of (a).
- 12. (PREVIOUSLY AMENDED) The set of oligonucleotides of claim 11, wherein said set consists of forward primers and reverse primers and hybridization probes.

- 13. (CURRENTLY AMENDED) The set of oligonucleotides of claim 11, wherein each oligonucleotide eomprises-consists of one of SEQ ID NOS: 1, 2, 3, 5, 6, and 7.
- 14. (CURRENTLY AMENDED) A set of oligonucleotides comprising (a) a polynucleotide fragment of each of the isolated polynucleotides of the composition of claim 10, wherein said fragments are 12 to 50 19 to 33 nucleotides in length, or (b) full-length complements of (a).
- 15. (CURRENTLY AMENDED) The set of oligonucleotides of claim 14, wherein each oligonucleotide emprises-consists of one of SEQ ID NOS: 1, 2, 3, 5, 6, 7, 9, 10, 11, 13, 14, 15, 17, 18, 19, 21, 22, and 23.

16. (CANCELED)

17. (CANCELED)